

Fig. 1

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 CAGGTTCTCTCTTCAAATCATACACTTGGTTGAGTGAGAAATATGGACGTCAACCTAGCAATATGAATCCCTCTCCAAGATCTACACT -1320
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12bp element

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TATA box

transcriptional start site

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 AGCGAGACAACTCTATTAAACACCGGTAATTAACAACCGTTGTTGTCGAGTTCATGTTTCTTCCAACTCTTTCTCTTTCTTACTTT 121
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1

M N Q E M

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 ATATTGTCGAGGTTACAAAGAAATGAGTCTGTTTGGACTCAAAATCAACATTGGTGTGAGTTTCAAACTAAACTGTCATTAAT 391
 36 I L S R F T K N E F C F D S K S T I G V E F Q T K T V T I N

GGTAACTCATCAAGCTCAGATCTGGGACTGCTGGCAAGAAAGGTTTCTTCTCTTCTCTCTATAATCTCGAATATATATTTTACT 481
 66 G K L I K A Q I W D T A G Q E R
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133 N K G D L V D Q R G V Q T E D A V E F A E D Q G L F F S E T

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163 S A F S G E N V N S A F L K L L Q E I N K V V S K R S L E C

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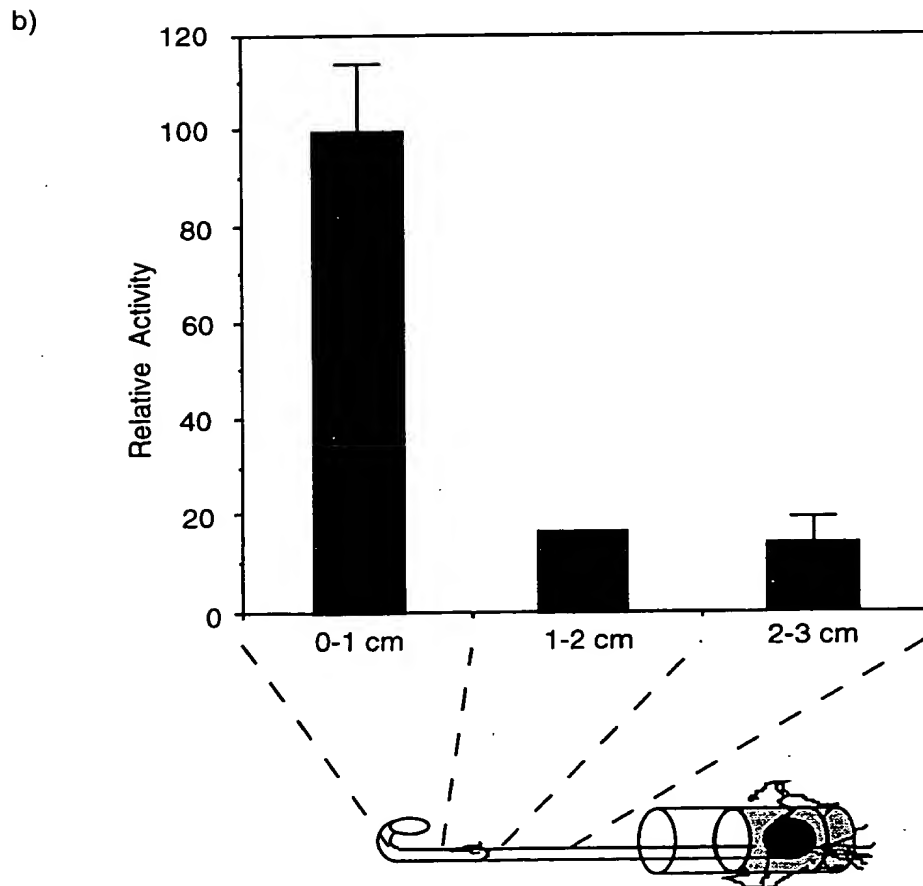
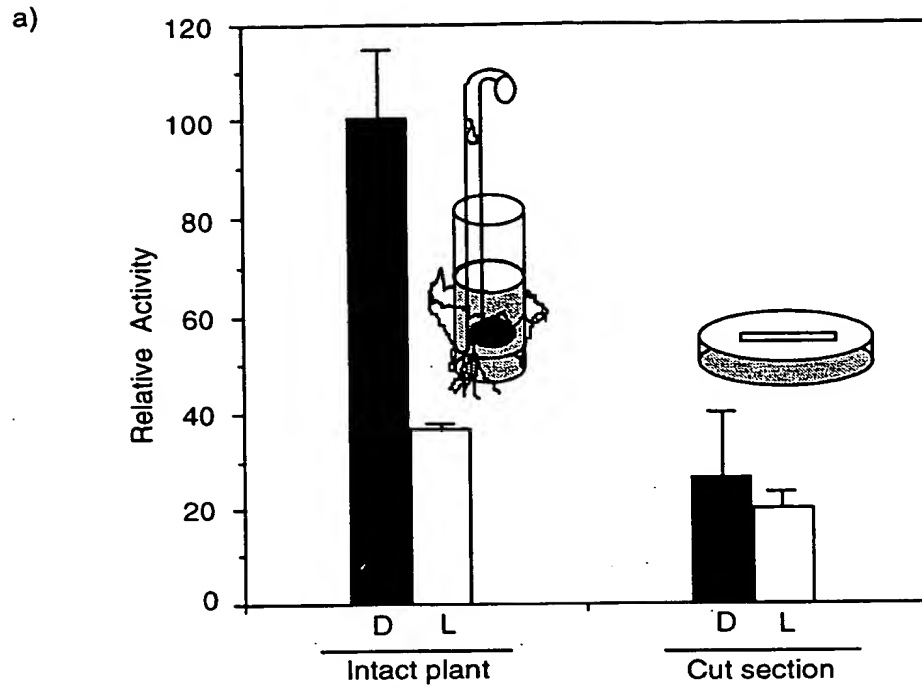
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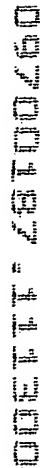
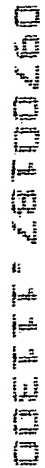
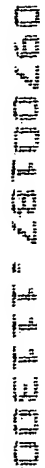
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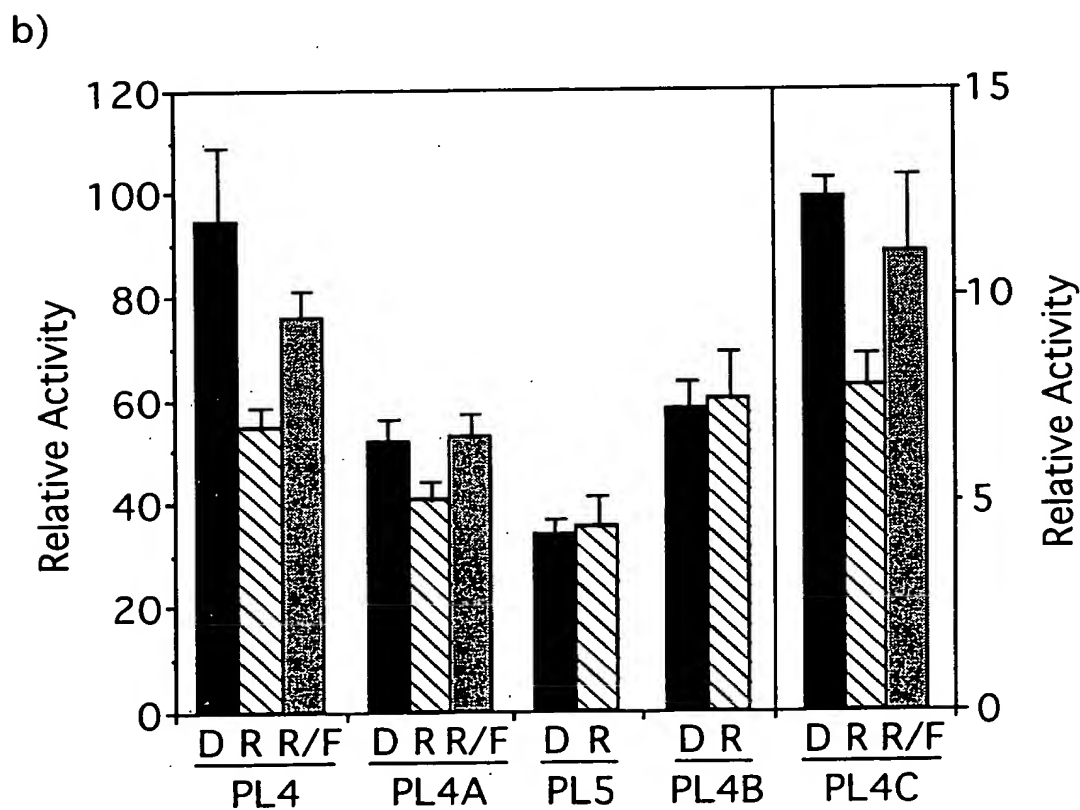
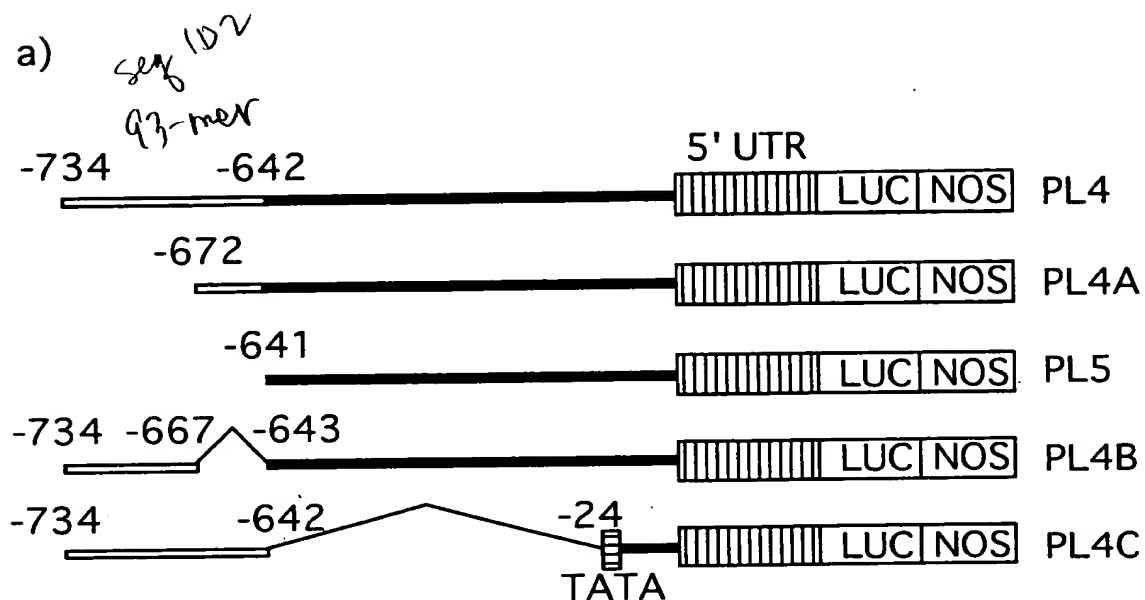
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Fig. 2







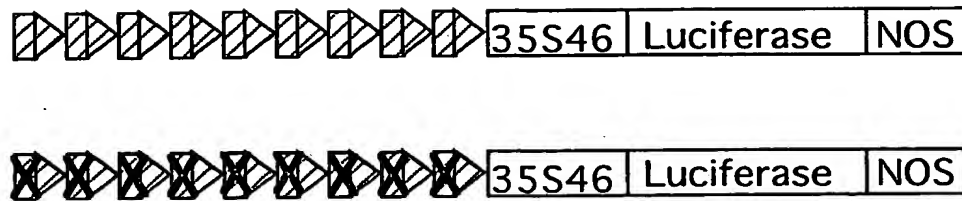
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Relative Activity

Group	D (Solid Bar)	R (Hatched Bar)
LS1	40.5	27.5
LS2	34.5	29.5
LS3	26.0	24.0
LS4	33.5	25.5
LS5	38.5	28.5

Fig. 8

a)



b)

